SEQUENCE LISTING

Armour, Christopher Castle, John Garrett-Engele, Philip Johnson, Jason

<120> ALTERNATIVELY SPLICED ISOFORMS OF HUMAN PHKA2

<130> RS0204Y

<140> US 10/648,139

<141> 2003-08-26

<150> US 60/408,058

<151> 2002-09-03

<150> US 60/431,474

<151> 2002-12-05

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Leu Ser Ala Ser His Glu Gln Lys Asp Ala Trp Val Arg Asp Asn Ile 35 40 45

Tyr Ser Ile Leu Ala Val Trp Gly Leu Gly Met Ala Tyr Arg Lys Asn 50 60

Ala Asp Arg Asp Glu Asp Lys Ala Lys Ala Tyr Glu Leu Glu Gln Asn 65 70 75 80

Val Val Lys Leu Met Arg Gly Leu Leu Gln Cys Met Met Arg Gln Val 85 90 95

Ala Lys Val Glu Lys Phe Lys His Thr Gln Ser Thr Lys Asp Ser Leu 100 105 110

His Ala Lys Tyr Asn Thr Ala Thr Cys Gly Thr Val Val Gly Asp Asp 115 120 125

Gln Trp Gly His Leu Gln Val Asp Ala Thr Ser Leu Phe Leu Leu Phe 130 135 140

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Lys Val Ala Asp Tyr Gly Met Trp Glu Arg Gly Asp Lys Thr Asn Gln 180 185 190

Gly Ile Pro Glu Leu Asn Ala Ser Ser Val Gly Met Ala Lys Ala Ala 195 200 205

Leu Glu Ala Ile Asp Glu Leu Asp Leu Phe Gly Ala His Gly Gly Arg 210 215 220

Lys Ser Val Ile His Val Leu Pro Asp Glu Val Glu His Cys Gln Ser 225 230 235 240

Ile Leu Phe Ser Met Leu Pro Arg Ala Ser Thr Ser Lys Glu Ile Asp 245 250 255

Ala Gly Leu Leu Ser Ile Ile Ser Phe Pro Ala Phe Ala Val Glu Asp 260 265 270

Val Asn Leu Val Asn Val Thr Lys Asn Glu Ile Ile Ser Lys Leu Gln 275 280 285

Gly Arg Tyr Gly Cys Cys Arg Phe Leu Arg Asp Gly Tyr Lys Thr Pro 290 295 300

Arg Glu Asp Pro Asn Arg Leu His Tyr Asp Pro Ala Glu Leu Lys Leu 305 310 315 320

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Ile Asp Gly Val Phe Ser Gly Asp Ala Val Gln Val Gln Glu Tyr Arg 340 345 350

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Asp Ile His Ser Ala Val Leu Ser Thr Ile Arg Lys Leu Glu Asp Gly Tyr Phe Gly Gly Ala Arg Val Lys Leu Gly Asn Leu Ser Glu Phe Leu Thr Thr Ser Phe Tyr Thr Tyr Leu Thr Phe Leu Asp Pro Asp Cys Asp Glu Lys Leu Phe Asp Asn Ala Ser Glu Gly Thr Phe Ser Pro Asp Ser Asp Ser Asp Leu Val Gly Tyr Leu Glu Asp Thr Cys Asn Gln Glu Ser Gln Asp Glu Leu Asp His Tyr Ile Asn His Leu Leu Gln Ser Thr Ser Leu Arg Ser Tyr Leu Pro Pro Leu Cys Lys Asn Thr Glu Asp Arg His Val Phe Ser Ala Ile His Ser Thr Arg Asp Ile Leu Ser Val Met Ala Lys Ala Lys Gly Leu Glu Val Pro Phe Val Pro Met Thr Leu Pro Thr Lys Val Leu Ser Ala His Arg Lys Ser Leu Asn Leu Val Asp Ser Pro Gln Pro Leu Glu Lys Val Pro Glu Ser Asp Phe Gln Trp Pro Arg Asp Asp His Gly Asp Val Asp Cys Glu Lys Leu Val Glu Gln Leu Lys Asp Cys Ser Asn Leu Gln Asp Gln Ala Asp Ile Leu Tyr Ile Leu Tyr Val Ile Lys Gly Pro Ser Trp Asp Thr Asn Leu Ser Gly Gln His Gly

Val Thr Val Gln Asn Leu Leu Gly Glu Leu Tyr Gly Lys Ala Gly Leu 805 810 815 Asn Gln Glu Trp Gly Leu Ile Arg Tyr Ile Ser Gly Leu Leu Arg Lys 820 825 Lys Val Glu Val Leu Ala Glu Ala Cys Thr Asp Leu Leu Ser His Gln Lys Gln Leu Thr Val Gly Leu Pro Pro Glu Pro Arg Glu Lys Ile Ile 855 860 Ser Ala Pro Leu Pro Pro Glu Glu Leu Thr Lys Leu Ile Tyr Glu Ala 865 870 875 Ser Gly Gln Asp Ile Ser Ile Ala Val Leu Thr Gln Glu Ile Val Val 885 890 895 Tyr Leu Ala Met Tyr Val Arg Ala Gln Pro Ser Leu Phe Val Glu Met 900 905 910 Leu Arg Leu Arg Ile Gly Leu Ile Ile Gln Val Met Ala Thr Glu Leu 915 920 Ala Arg Ser Leu Asn Cys Ser Gly Glu Glu Ala Ser Glu Ser Leu Met 935 Asn Leu Ser Pro Phe Asp Met Lys Asn Leu Leu His His Ile Leu Ser 955 Gly Lys Glu Phe Gly Val Glu Arg Ser Glu Val Gln His Pro Ile Leu 965 970

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Ala Asp Arg Asp Glu Asp Lys Ala Lys Ala Tyr Glu Leu Glu Gln Asn 65 70 75 80

Val Val Lys Leu Met Arg Gly Leu Leu Gln Cys Met Met Arg Gln Val 85 90 95

Ala Lys Val Glu Lys Phe Lys His Thr Gln Ser Thr Lys Asp Ser Leu 100 105 110

His Ala Lys Tyr Asn Thr Ala Thr Cys Gly Thr Val Val Gly Asp Asp 115 120 125

Gln Trp Gly His Leu Gln Val Asp Ala Thr Ser Leu Phe Leu Leu Phe 130 135 140

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Ser Val Lys Pro Asp Val Val Val Gln Val Thr Val Leu Ala Glu Asn 435 440 Asn His Ile Lys Asp Leu Leu Arg Lys His Gly Val Asn Val Gln Ser Ile Ala Asp Ile His Pro Ile Gln Val Gln Pro Gly Arg Ile Leu Ser His Ile Tyr Ala Lys Leu Gly Arg Asn Lys Asn Met Asn Leu Ser Gly 490 Arg Pro Tyr Arg His Ile Gly Val Leu Gly Thr Ser Lys Leu Tyr Val 500 505 Ile Arg Asn Gln Ile Phe Thr Phe Thr Pro Gln Gln Met Met Ala Gln 515 520 525 Thr Phe Ile Leu Leu Cys Ser Pro Gln Leu Glu Asn 530 535 <210> 5 1779 <211> <212> DNA <213> Homo sapiens <400> atgcggagca ggagcaattc cggggtccgc ttggacgggt acgcgcggct ggtgcagcaa 60 accatectgt gttaccagaa teeegteacg gggetgetgt cagecageca tgageagaag 120 gatgcctggg tgcgggataa catctacagt atcctggccg tgtggggcct gggcatggcc 180 taccgtaaga atgcagaccg cgatgaggac aaggccaagg cctacgagct ggagcagaac gtggtgaage tgatgegagg tetteteeag tgeatgatga gaeaggtgge caaagtggag 300 aagttcaaac acactcagag caccaaggac agcctgcacg ccaagtacaa caccgccacc 360 tgtggcacgg tggtgggcga cgaccagtgg ggccacctcc aggtggatgc cacctctctc 420 ttcctcctgt tcctggccca gatgaccgcc tcaggcttac gtatcatttt cactctcgat 480 gaggtggcct tcatacagaa tcttgtcttt tacatagaag ctgcatataa agtcgctgat 540 tatggaatgt gggagcgtgg agataagact aatcagggca tcccggaatt gaatgcaagc 600

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720

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Ile 465	Ala	Asp	Ile	His	Pro 470	Ile	Gln	Val	Gln	Pro 475	Gly	Arg	Ile	Leu	Ser 480
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